

Specific binding of *Clostridium perfringens* enterotoxin fragment to Claudin-b and modulation of zebrafish epidermal barrier

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Abstract: Claudins (Cldn) are the major components of tight junctions (TJs) sealing the paracellular cleft in tissue barriers of various organs. Zebrafish Cldnb, the homolog of mammalian Cldn4, is expressed at epithelial cell-cell contacts and is important for regulating epidermal permeability. The bacterial toxin *Clostridium perfringens* enterotoxin (CPE) has been shown to bind to a subset of mammalian Cldns. In this study, we used the Cldn-binding C-terminal domain of CPE (194–319 amino acids, cCPE_{194–319}) to investigate its functional role in modulating zebrafish larval epidermal barriers. *In vitro* analyses show that cCPE_{194–319} removed Cldn4 from epithelial cells and disrupted the monolayer tightness, which could be rescued by the removal of cCPE_{194–319}. Incubation of zebrafish larvae with cCPE_{194–319} removed Cldnb specifically from the epidermal cell membrane.

Dye diffusion analysis with 4-kDa fluorescent dextran indicated that the permeability of the epidermal barrier increased due to cCPE_{194–319} incubation. Electron microscopic investigation revealed reversible loss of TJ integrity by Cldnb removal. Collectively, these results suggest that cCPE_{194–319} could be used as a Cldnb modulator to transiently open the epidermal barrier in zebrafish. In addition, zebrafish might be used as an *in vivo* system to investigate the capability of cCPE to enhance drug delivery across tissue barriers.

Key words: Claudin-4 – Claudin-b – *Clostridium perfringens* enterotoxin – epidermal barrier – tight junction

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Introduction

Epithelial barriers cover the surfaces of multicellular organisms, insulating diverse tissue compartments from the external environment and maintaining homeostasis (1). In vertebrates, epithelial tissues include gastrointestinal and respiratory epithelia, epidermal squamous epithelium, accessory glands, retinal pigment epithelium of the eye and renal epithelium (2,3). The exchange of substances through intercellular spaces between external environment and tissues is constrained by the epithelial barrier. Two routes across epithelia are generally adopted for drug delivery: the transcellular pathway and paracellular pathway (3,4). However, in the later, therapeutic drug delivery is often retarded by the epithelial barrier. Strategies that transiently break down the epithelial barrier and enhance drug transport are in need.

Tight junctions (TJs) organized in strand networks at the cell-cell contacts between epithelial cells are the molecular structural basis of the epithelial barrier (4). TJs span the plasma membrane of epithelial cells at the interface between the apical and basolateral parts. By interactions between TJ strands of neighbouring cells, TJs form a barrier in the extracellular space. Consequently, these cells together with their TJs establish a barrier which directly regulates paracellular flux (5).

Tight junction complexes are mainly composed of the transmembrane proteins occludin, Claudins (Cldns) and various scaffolding proteins, such as zonula occludens proteins -1, -2 and

-3 (ZO-1, -2 and -3) (5). Cldns play a dominant role on the properties of the TJs (6–9). They are expressed in a tissue-specific combination manner, resulting in tissue-specific barrier function. Targeting Cldns presents a potential solution to open TJs and enhance drug delivery (3,4).

At present, a variety of strategies have been designed to breach TJs to improve the paracellular passage of drugs through epithelial barriers. Shorter peptides homologous to external residues of TJ proteins, specific targeting TJ proteins by siRNA methods, and bacterial toxins causing disruption of TJs are vigorously explored for this purpose (3). Recently, an enterotoxin derived from *Clostridium perfringens* (CPE) was found to bind to Cldns 3, 4, 6, 7, 8 and 14 (10,11). CPE could disintegrate TJs and induce permeability increase of epithelial barrier. The C-terminal portion of CPE (cCPE), which lacks cytotoxic activity and has the Cldn-binding domain, could remove specific Cldns from TJs and induce opening of TJs (12,13). Short peptides derived from CPE could possibly be used to promote transepithelial drug delivery (3).

In this study, we apply a truncated CPE fragment containing amino acids from 194 to 319 (cCPE_{194–319}) to investigate its functional role in modulating zebrafish larval epidermal barriers. Zebrafish Cldnb, the homolog of mammalian Cldn4, is expressed at developing zebrafish epidermal cell-cell contacts (14). The results showed that cCPE_{194–319} bonded and cleared Cldn4 from TJs in mammalian epithelial cells. The cCPE_{194–319} also conjugated

with *Cldnb*, induced severe pericardial oedema, and boosted the diffusion of 4-kDa fluorescent dextran (FD-4 kDa) across epidermis in zebrafish. Collectively, these results suggest that *cCPE*_{194–319} could modulate transient breakdown of epidermal barrier by regulating *Cldnb* and possibly be used as a transepithelial enhancer during drug delivery.

Materials and methods

Zebrafish husbandry

Adult AB zebrafish were maintained and raised at 28.5°C under standard conditions. Embryos were kept in egg water according to standard method (15). The stage of the embryos was determined by morphological features and by hours postfertilization (hpf) at 28.5°C (16). 0.003% (w/v) 1-phenyl-2-thiourea (PTU; Sigma-Aldrich, St. Louis, MO, USA) was added into the egg water to suppress pigmentation. Handling of zebrafish and mice was in compliance with 'The legislation of Guangdong laboratory animal management regulations', carefully monitored and approved by the ethic committee of the Affiliated Hospital, Guangdong Medical College.

Protein synthesis and purification

For glutathione-S-transferase (GST) and GST-fused C-terminal of *C. perfringens* enterotoxin (GST-*cCPE*_{194–319}) purification, glycerol stocks of transformed *Escherichia coli* BL21 (DE3) were cultured and purified using glutathione-sepharose 4B beads (Amersham, Buckinghamshire, UK). Eluted GST fusion proteins were dialysed against PBS at 4°C and stored for short time at 4°C and prolonged time at –80°C for further experiments.

Cell culture, immunocytochemistry and trans-epithelial electric resistance measurement

MDCK-I cells were grown on poly-L-Lysine (Sigma-Aldrich)-coated coverslips at 10% CO₂ in DMEM supplemented with 10% FBS (Gibco, Grand Island, NY, USA). Cells were plated in a density of 100 000 cells/cm² and grown for 48 h until it formed a confluent monolayer. For *cCPE*_{194–319} treatment, purified GST-*cCPE*_{194–319} was added into culture medium at a final concentration of 2 µg/ml. Cells were incubated for 30 min and 24 h, respectively. For the recovery experiments, *cCPE*_{194–319}-containing medium was removed after 24-h incubation. Cells were washed with PBS for 5 min and incubated with fresh culture medium for another 24 h. For immunocytochemical staining, cells were fixed and stained with rabbit anti-GST or mouse anti-*Cldn4* antibodies. DAPI (4',6-diamidino-2-phenylindole dihydrochloride, 1:1000; Molecular Probes, Grand Island, NY, USA), goat anti-rabbit Cy3 (1:200; Jackson ImmunoResearch, West Grove, PA, USA) and goat anti-mouse FITC (1:200; Jackson ImmunoResearch) were used as secondary antibodies. For confocal imaging, a Leica TSC SP5II (Leica, Solms, Germany) confocal microscope and a 60× objective was used. Trans-epithelial electric resistance (TEER) measurements were performed using EVOM² epithelial voltohmmeter (WPI, Sarasota, FL, USA). When MDCK-I cells were cultured on rat tail collagen precoated Transwell™ chamber (Millipore, USA), TEER was measured at series time points under different conditions. For each condition, 18 filters with MDCK-I cells were investigated.

Morpholino injection and *cCPE*_{194–319} incubation

Zebrafish *cldnb* morpholino was used as described (5'-CCG GTT GAT GCC ATG CTT TTT CGT T-3') (Genetools, Philomath, OR, USA). Morpholino was injected at one cell stage at a concentration of 200 µM. For GST-*cCPE*_{194–319} incubation, 24-hpf embryos

were dechlorinated and incubated in a 24-well plate with 50 µg/ml GST-*cCPE*_{194–319} in egg water. After 24- or 48-h incubation, fresh egg water was changed and embryos were incubated for another 24 h. For imaging, embryos were mounted in 1% low melting agarose (LMA; Invitrogen, Grand Island, NY, USA). Leica M205FA stereo microscope was used to take differential interference contrast images. Heart beating was counted with live embryos under the stereo microscope. Twenty-seven embryos were analysed for each group.

Embryo whole-mount immunohistochemistry

Embryos were washed with PBS three times for 5 min each and fixed in 2% paraformaldehyde (PFA; Sigma-Aldrich) at 4°C overnight. For immunohistochemical staining, the following antibodies were used: mouse anti-*Cldn4* which also recognizes zebrafish *Cldnb* (1:100; Invitrogen), rabbit anti-ZO-1 (1:200; Invitrogen), rabbit anti-GST (1:200; Santa Cruz, Dallas, Texas, USA), goat anti-rabbit Cy5 (1:200; Jackson ImmunoResearch Laboratories), goat anti-mouse FITC (1:100; Jackson ImmunoResearch Laboratories), rhodamine phalloidin (1:500; Invitrogen), 4',6-Diamidino-2-phenylindole dihydrochloride (1:1000; Sigma-Aldrich). Confocal images were obtained with a Leica TSC SP5II (Leica) confocal microscope. Images were processed using Photoshop software (Adobe, San Jose, CA, USA).

Zebrafish epidermal barrier dye diffusion assay

Embryos were incubated with 50 µg/ml 4-kDa fluorescent dextran (Sigma-Aldrich) contained egg water for 30 min covered from light, followed by being rinsed with egg water for three times 5 min each. Embryos were then mounted in 1% LMA and imaged with Leica TSC SP5II confocal microscope (Leica). Images were processed using Photoshop software (Adobe).

Transmission electron microscopy analysis

Embryos were fixed with 2.5% glutaraldehyde (Sigma-Aldrich) buffered in 0.1 M cacodylate buffer (pH 7.4). Thereafter, the whole fish was postfixed in 1% OsO₄ in 0.1 M cacodylate buffer and dehydrated in graded ethanol. The specimens were then embedded in plastics (Araldite based). Thin sections were made from plastic blocks on a FCR Reichert Ultracut ultramicrotome (Leica) and stained with uranyl acetate/lead citrate. Section slices were mounted on pioloform-coated copper grids, and images were analysed and documented with a JEM-1400 electron microscope (JEM, Tokyo, Japan). Seven samples of each group were analysed.

Mice epidermal barrier investigation assay

Adult BALB/c mice were purchased from the Animal Center of Guangdong Medical College. Firstly, GST-*cCPE*_{194–319} or GST constructs were applied on the skin of living mice. Then, after their sacrifice, diffusion of fluorescent dye was analysed. In brief, the hair was first cut off by hair cut and then removed by hair removal cream carefully, before tape stripping to remove the stratum corneum. For *cCPE*_{194–319} incubation, living mice were anaesthetized for 30 min with 2% pentobarbital sodium (40 mg/kg; Merck, Kenilworth, NJ, Germany). Then, 100 µg/ml *cCPE*_{194–319} or GST was added on a wound plaster and stuck onto the nude skin of the mice back. Wound plaster was re-fixed with a 3D self-adhesive elastic bandage to prevent the dropping off of the plaster. After 24-h treatment, mice were sacrificed by cervical dislocation and the skin regions incubated with the recombinant protein were cut off and washed with PBS. Skins were incubated with 10 µg/ml 10-kDa rhodamine B dextran (Sigma-Aldrich) for 10 min

protected from light, then washed with PBS for 6×5 min each. For imaging, skin blocks were mounted in slowfade[®] diamond antifade mountant (Life Technologies, Grand Island, NY, USA) and imaged with Leica M205 FA stereomicroscope (Leica). To ensure the stratum corneum was clearly removed after the tape stripping, the paraffin section of the same skin block and the haematoxylin-eosin staining was then performed and imaged with Leica DM4000 B microscope (Leica). Six mice for each condition (GST-cCPE₁₉₄₋₃₁₉ or GST alone, as a control) were analysed.

Results

The C-terminal domain of *Clostridium perfringens* enterotoxin (cCPE₁₉₄₋₃₁₉) binds to Claudin-4 and modulates paracellular barrier in cell monolayers

C-terminal fragments of the *C. perfringens* enterotoxin have been reported to selectively bind to and to remove Cldn4 from TJs (12, 17). In this study, we used recombinant GST-cCPE₁₉₄₋₃₁₉ fusion protein to bind to Cldns and thereby to modulate TJs in order to investigate the regulation of tissue barriers. First, the activity of GST-cCPE₁₉₄₋₃₁₉ was analysed employing MDCK-I cells as an established cell culture model system for epithelial barriers. After 30-min incubation of MDCK-I cells monolayer, GST-cCPE₁₉₄₋₃₁₉, but not the GST control, was found to bind strongly to the cell membrane, mainly at cell-cell contacts where Cldn4 was present (Fig. 1a-i). After 24 h of incubation with GST-cCPE₁₉₄₋₃₁₉, Cldn4 almost completely disappeared from the cell membrane (Fig. 1j-o). Although GST-cCPE₁₉₄₋₃₁₉ was still strongly present on the cell membrane at this time, it could be easily washed away (Fig. 1j-l,p-r). After recovery for 24 h, Cldn4 reappeared at cell-cell contacts (Fig. 1p-r).

To study the effect of cCPE₁₉₄₋₃₁₉ on the epithelial barrier directly, the transepithelial electrical resistance (TEER) as a measure of the tightness of MDCK-I cell monolayers was measured. As shown in Fig. 2, MDCK-I cells formed a stable monolayer barrier 120 h after plating. When GST-cCPE₁₉₄₋₃₁₉ was added onto the monolayer at 144 h, TEER dropped from 1500 to 200 $\Omega\cdot\text{cm}^2$ within the following 24 h. In contrast, GST applied as a control to the cell monolayer showed no decrease of TEER. To test whether the leaky epithelial monolayer can recover tightness, GST-cCPE₁₉₄₋₃₁₉ was removed at 168 h. Removal of GST-cCPE₁₉₄₋₃₁₉ induced the recovery of TEER, which reached the level of 1200 $\Omega\cdot\text{cm}^2$ within 24 h. Taken together, these results demonstrate that the expressed and purified GST-cCPE₁₉₄₋₃₁₉ is functionally active and able to reversibly increase the paracellular permeability for ions by modulating the barrier-forming TJ protein, Cldn4.

cCPE₁₉₄₋₃₁₉ dislocates Cldnb in zebrafish embryos

We aimed to explore whether cCPE₁₉₄₋₃₁₉ could be used to temporarily modulate the zebrafish epidermal barrier *in vivo*. Zebrafish Cldnb is orthologous to mammalian Cldn4 (14). The turn region in the ECL2 of Cldnb (149NPLVV153) is very similar to the cCPE binding motif (NPL/MVA/P) in mammalian Cldns (18). However, it is unknown whether cCPE₁₉₄₋₃₁₉ is able to interact with Cldnb. To answer this question, zebrafish larvae were incubated with GST-cCPE₁₉₄₋₃₁₉ and immunohistochemical analysis of the zebrafish trunk epidermis was performed. Without treatment, Cldnb and TJ marker ZO-1 were found to be expressed and to colocalize together on the epithelial cell-cell contacts at 48 hpf (Fig. 3a-c). When zebrafish embryos (24 hpf) were incubated with GST-cCPE₁₉₄₋₃₁₉ for 3 h, GST-cCPE₁₉₄₋₃₁₉ colocalized with Cldnb

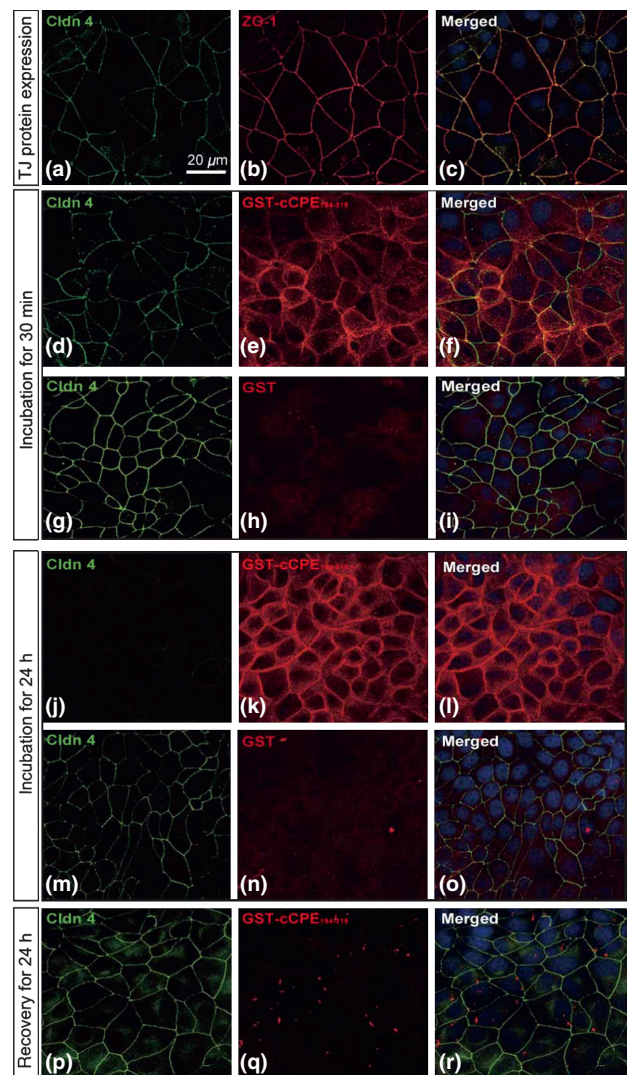


Figure 1. C-terminal fragment of *Clostridium perfringens* enterotoxin (cCPE₁₉₄₋₃₁₉) binds to MDCK-I cells and removes Cldn4 from tight junctions (TJs). Immunocytochemical staining and confocal microscopic images. (a-c) The TJ components Cldn4 and ZO-1 colocalized at the cell-cell contact of confluent MDCK-I cells. (d-f) GST-cCPE₁₉₄₋₃₁₉ (red) colocalized with Cldn4 (green) on the MDCK-I cell membrane after 30 min of incubation. (g-i) As a control, binding of GST (red) alone to the cells was not detected. (j-l) Specific binding of GST-cCPE₁₉₄₋₃₁₉ (red) causes loss of Cldn4 (green) from MDCK-I cell-cell contacts after 24-h incubation. (m-o) In contrast, GST (red) incubation has no effect on the presents of Cldn4 (green) at cell-cell contacts. (p-r) After 24-h incubation, GST-cCPE₁₉₄₋₃₁₉ was washed away and the MDCK-I cells were incubated with fresh cell culture media for another 24 h. Cldn4 (green) and reappeared at cell-cell contacts of MDCK-I cells. Scale bar, 20 μm . GST, glutathione-S-transferase.

on the cell membrane (Figure S2a-d). However, when Cldnb was knocked down, GST-cCPE₁₉₄₋₃₁₉ was not able to bind to the epidermal cells which were visualized by F-actin staining. This finding indicated that Cldnb is necessary for the binding of GST-cCPE₁₉₄₋₃₁₉ to the epidermal cells (Figure S2e-l). After 24 h of incubation, Cldnb disappeared from the epidermal cell membrane, without affecting the localization of ZO-1 (Fig. 3g-i), which was reminiscent of the effect of specific morpholino oligonucleotide targeting Cldnb (Fig. 3d-f). These results suggest that cCPE₁₉₄₋₃₁₉ is able to bind and modulate Cldnb.

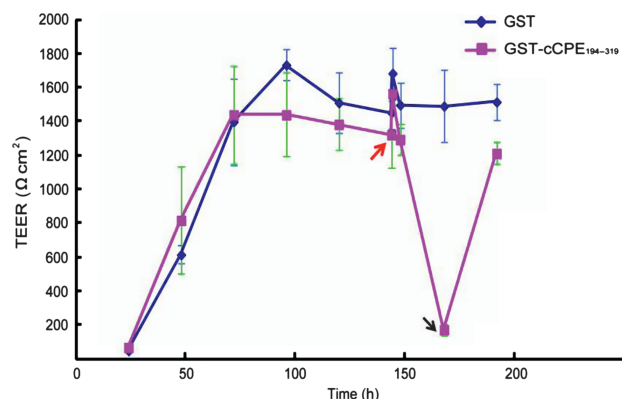


Figure 2. GST-cCPE₁₉₄₋₃₁₉ incubation reduces transepithelial electric resistance (TEER) of MDCK-I cell monolayer measuring. MDCK-I cells cultured for 120 h form a stable tight monolayer which reaches about 1500 Ω·cm² evaluated by measuring TEER. Application of 50 μg/ml GST-cCPE₁₉₄₋₃₁₉ (pink line) at 144 h (red arrow) reduced TEER down to about 200 Ω·cm² within the following 24 h. This indicates the increase of permeability for paracellular ion and the breakdown of paracellular barrier. In contrast, incubation of 50 μg/ml GST (blue line) did not decrease TEER. Washout of GST-cCPE₁₉₄₋₃₁₉ by replacement with fresh media (black arrow) at 168 h leads to the recovery of TEER back to 1200 Ω·cm² within the following 24-h incubation. CPE, *Clostridium perfringens* enterotoxin; GST, glutathione-S-transferase.

cCPE₁₉₄₋₃₁₉ causes transitory pericardial oedema in zebrafish embryos

The deficiency of Cldnb has been reported to cause developmental defects in zebrafish (14). In accordance with this prior report, knock-down of Cldnb by morpholino oligonucleotide elicited dramatic pericardial oedema and curly tail in the zebrafish morphants (Figure S1a–d). However, elimination of Cldnb by GST-cCPE₁₉₄₋₃₁₉ only induced similar oedema in the pericardial cavity but not curly tail phenotype, when zebrafish embryos were incubated with GST-cCPE₁₉₄₋₃₁₉ for 24 h (Figure S1e,f). The extension of incubation time with GST-cCPE₁₉₄₋₃₁₉ to 48 h aggravated the oedema without affecting the tail (Figure S3a–f). Interestingly, when the affected zebrafish embryos were taken back into fresh egg water, pericardial oedema recovered autonomously (Figure S3 g,h). In addition, the cCPE₁₉₄₋₃₁₉ incubation severely reduced the heart beating rate in the zebrafish, which could recover after the clearance of cCPE₁₉₄₋₃₁₉ (Table S1). These results indicate that cCPE₁₉₄₋₃₁₉ could induce transitory disruption of epithelial barrier of the pericardial cavity in zebrafish embryos.

cCPE₁₉₄₋₃₁₉ temporarily boosts penetration of 4-kDa dextran through zebrafish epidermal barrier

To examine whether cCPE₁₉₄₋₃₁₉ could facilitate the penetration of drugs through epithelial barriers, the permeability of the zebrafish epidermis for fluorescent dextran was studied. As shown in Fig. 4a,b, zebrafish epidermal cells formed a tight epithelial barrier at 48 hpf which is not permeable to 4-kDa FITC-dextran (FD-4 kDa). Transmission electron microscopy of zebrafish larval epidermis also revealed intact TJ (red arrow) at the apical side of adjacent epithelial cell–cell contacts (Fig. 4c). In contrast, removal of Cldnb by incubation of zebrafish larvae with morpholino oligonucleotides (Fig. 4d,e) or with GST-cCPE₁₉₄₋₃₁₉ (Fig. 4g,h) increased the penetration of FD-4 kDa through the epidermis. In line with the increased dextran permeability for these two groups, disruption of TJs between the neighbouring epithelial cells was observed on the ultrastructural level (Fig. 4f,i). However, when

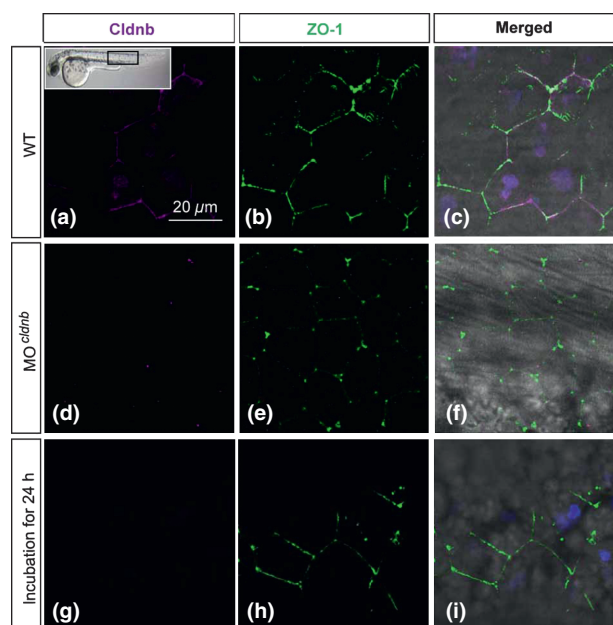


Figure 3. Incubation of zebrafish larvae with GST-cCPE₁₉₄₋₃₁₉ contained egg water causes loss of Cldnb on zebrafish epidermal cell membrane. (a–c) Immunohistochemical staining and confocal microscopy on the trunk epidermis shows that Cldnb and ZO-1 are expressed and colocalize on the zebrafish skin epithelial cell–cell contacts at 48 hpf. (d–f) Specific knock-down of *cldnb* by morpholino oligonucleotide reduces the expression of Cldnb on the epithelial cell membrane, while the expression of ZO-1 is not affected. (g–i) Incubation of 24-hpf zebrafish embryos with 50 μg/ml GST-cCPE₁₉₄₋₃₁₉ proteins for 24 h causes the disappearance of Cldnb from epidermis cell membrane without affecting the localization of ZO-1. Scale bar, 20 μm. CPE, *Clostridium perfringens* enterotoxin; GST, glutathione-S-transferase.

GST-cCPE₁₉₄₋₃₁₉-treated zebrafish embryos were incubated with fresh egg water for another 24 h, embryos displayed low permeability for FD-4 kDa, indicating recovery of the TJ and epithelial barrier (Fig. 4j–l). In summary, these results show that cCPE₁₉₄₋₃₁₉ can reversibly enhance the penetration of molecules at least up to 4 kDa through the zebrafish epidermal barrier.

To ask whether cCPE₁₉₄₋₃₁₉ could also promote drug delivery through mammalian epidermal barriers, we also performed preliminary analyses on mouse skin after removing the stratum corneum by tape stripping. As a result, cCPE₁₉₄₋₃₁₉ treatment improved the diffusion of rhodamine B dextran (10 kDa) through the mice epidermal barrier (Figure S4).

Discussion

This study uses a zebrafish model to test the *in vivo* capability of recombinant cCPE₁₉₄₋₃₁₉ to act as a Cldn modulator for analysis of epithelial barrier functions, or for enhancement of paracellular drug delivery across tissue barriers. The results confirm that cCPE₁₉₄₋₃₁₉ reversibly binds to and removes Cldn4 present on the plasma membrane of mammalian epithelial cells *in vitro*. Further experiments show that cCPE₁₉₄₋₃₁₉ incubation causes zebrafish temporal pericardial oedema due to breakdown of the epidermal barrier. The cCPE₁₉₄₋₃₁₉ conjuncts to zebrafish Cldnb and withdraws Cldnb from the epidermis. In accordance, cCPE₁₉₄₋₃₁₉ improves the permeability for FD-4 kDa through the zebrafish epidermis. Importantly, the cCPE₁₉₄₋₃₁₉ induced opening of epithelial barrier is reversible, leaving no apparent harm after barrier

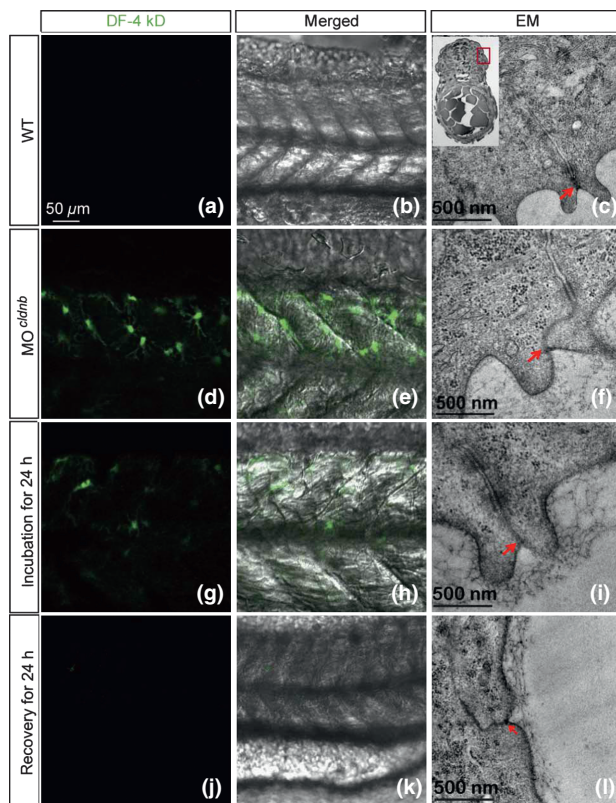


Figure 4. Loss of Cldnb on epidermis cell membrane breaks down the epithelial tight junction (TJ) and increases the skin epithelial permeability for 4-kDa dextran in zebrafish larvae at 3 dpf (days postfertilization). (a, b) Zebrafish epidermis cells form a tight epithelium and epithelial barrier at 48 hpf which is not permeable to FD-4 kDa after 30-min incubation. (c) Transmission electron microscopy (TEM) analysis of zebrafish larva epidermis shows intact TJ strand (red arrow) at the apical side of adjacent epithelial cell–cell contact. Inset indicates the epithelial cell–cell contact side of the trunk region for EM analysis. (d, e, g and h) Both knock-down of *cldnb* by MO and incubation of zebrafish larvae with GST-cCPE_{194–319} proteins for 24 hpf increase permeability for FD-4 kDa through embryonic epidermis at 48 hpf. After 30-min incubation, FD-4 kDa diffuses through the skin epidermis indicating affected epidermal barrier in the treated embryos. (f, i) TEM images indicate knock-down of *cldnb* or incubation of embryos with GST-cCPE_{194–319} proteins contained egg water breaks down the TJ between the neighbouring epithelial cells. (j, k) Incubation of GST-cCPE_{194–319}-treated zebrafish embryos with fresh egg water for additional 24 h, resulted in low permeability for FD-4 kDa after 30-min dye incubation, indicating recovery of the epithelial barrier. TEM analysis of epithelial cell–cell contacts of these embryos at 72 hpf shows (partly) recovered TJ between epithelial cells (l). Scale bar, a, d, g and j, 20 μ m; c, f, i and l, 500 nm. CPE, *Clostridium perfringens* enterotoxin; GST, glutathione-S-transferase.

recovery. These findings show that zebrafish can be used as a model to explore cCPE_{194–319} as an enhancer of paracellular permeability of molecules up to 4 kDa for transepithelial drug delivery, such as trans-intestinal delivery or even trans-epidermal delivery.

cCPE_{194–319} is a better candidate tool to modulate epithelial barrier than other C-terminal fragments of CPE. First of all, the cCPE_{194–319} is shown to strongly interact with Cldns in a subtype-specific manner. The C-terminal region 290–319 of CPE is sufficient for its binding to Cldns (19). However, longer CPE fragments have stronger binding capacity (20). The crystal structure of cCPE_{194–319} shows that region 290–319 is only the core Cldn-binding domain formed by strands β 8 and β 9 (21). In contrast, the cCPE_{194–319} contains a nine-strand β -sandwich which forms the whole Cldn-binding domain (21). Second, the cCPE_{194–319}

specifically and reversibly regulates Cldn proteins. In MDCK-I cells, cCPE_{184–319} was reported to interact with Cldn4 but not with Cldn1 (12). In accordance, the results here show that cCPE_{194–319} specifically binds to and removes Cldn4 in MDCK-I cells as well as removes Cldnb, the homolog of Cldn4 in zebrafish. The molecular mechanism of the interaction between cCPE/CPE and mammalian Claudins was analysed in detail (18, 19, 22–24). The published data, the findings of this study and the sequence of the ECL2 of Cldnb especially in the turn region (149NPLVV153) which is very similar to the cCPE binding motif (NPL/MVA/P) make a direct interaction between Cldnb and cCPE very likely. Importantly, the reduction of Cldns by cCPE_{194–319} is reversible. Washing out of cCPE_{194–319} could restore the expression of Cldns. cCPE_{194–319} reversibly opens TJs most likely due to the dynamic assembly and disassembly of TJs (25, 26) and sequestration of Cldns by cCPE preventing their incorporation in TJ strands. Within the 24-h recovery time, new Cldn molecules can be synthesized and transported to the plasma membrane where they participate in the formation of TJ strands. In addition, after washing out, dissociation of cCPE from Cldns might also reduce Cldn sequestration. Finally, like other C-terminal fragments of CPE, the cCPE_{194–319} transiently breaches epithelial barrier without significant toxic effects. In a previous toxicological study, Suzuki et al. (27) also showed that intravenous administration (5 mg/kg) or nasal cCPE administration (2 mg/kg) did not increase indicative biochemical markers of liver and kidney injuries. Incubation of zebrafish with cCPE_{194–319} induced pericardial oedema due to disruption of epithelial barrier. However, the treated zebrafish recovered well after exchanging freshwater.

For the first time, we defined the phenotypic changes of zebrafish and aspects of the underlying molecular mechanism by cCPE_{194–319} treatment. Epithelial barrier is important for the development of zebrafish. When the TJ protein ZO-3 was silenced using morpholinos, the morphants showed a loss of the epidermal barrier (28). Moreover, pericardial oedema, loss of blood circulation and tail fin malformations were present in the zebrafish embryos (28). Kwong and Perry (14) reported that knock-down of Cldnb elicited zebrafish developmental defects, including oedema in the pericardial cavity and yolk sac, as well as curly tail. In this study, the incubation of zebrafish with cCPE_{194–319} only induced pericardial oedema without affecting the tail fin. The cCPE_{194–319} also reduced heart rate in the zebrafish. Interestingly, after termination of cCPE_{194–319} treatment, pericardial oedema ameliorated and the heart rate of the zebrafish climbed closely to a normal level. These phenotype changes are owing to the loss of Cldnb caused by cCPE_{194–319}. These results are also in agreement with the previous report that cCPE_{184–319} incubation causes development defects in mouse embryos via eliminating the expression of Cldn4 and Cldn6 (17).

Cldn-based TJs contribute to the barrier function of the mammalian skin (8, 9, 29, 30). The cCPE_{194–319} may be used as a transepithelial enhancer in several possible ways during drug delivery. Basically, the cCPE_{194–319} could bind to Cldns and transiently break down the epithelial barrier to enhance the penetration of unrelated molecules. Apparently, cCPE_{194–319} might be used in drug delivery through mucosal epithelial barrier in nasal cavity, pulmonary tract or intestinal lumen (2, 4, 31, 32). It has been demonstrated that cCPE_{194–319} promoted the nasal mucosal

absorption of 4-kDa dextran as well as jejunal absorption of both 4- and 150-kDa dextrans (33). The cCPE_{184–319} has also been reported to enhance the jejunal absorption of 4-kDa dextran over 400-fold compared with a clinically used absorption enhancer, sodium caprate (34). Moreover, cCPE_{194–319} and other cCPEs have been shown to increase nasal, jejunal and pulmonary absorption of pharmaceutical peptide human parathyroid hormone derivative, hPTH (1–14, 17–36). Another possibility is the usage of cCPE_{194–319} in transdermal drug delivery (36, 37). On one hand, the cCPE_{194–319} may disrupt TJs in sweat duct, hair follicle and sebaceous gland to promote drug delivery through shunt or appendageal route (38, 39). On the other hand, skin barrier in mammals is composed by stratum corneum as well as TJs in the stratum granulosum (8, 9, 30, 40). It is possible to use cCPE_{194–319} to promote transdermal drug delivery after removing the stratum corneum (36, 37). Actually, in our study, cCPE_{194–319} treatment caused the diffusion of 10-kDa rhodamine B dextran through the mouse skin barrier after stripping off the stratum corneum.

Because the disruption of epithelial barrier may cause certain disease (2), it is also crucial to cease transepithelial enhancer treatment and seal the opening of epithelial barrier as needed. In this study, we showed that while the cCPE_{194–319} increased the diffusion of FD-4 kDa in zebrafish, the breach of epithelial barrier induced by cCPE_{194–319} recovered well after termination of cCPE_{194–319} treatment.

In conclusion, the results herein suggest that the cCPE_{194–319} reversibly modulates epithelial barrier *in vivo* and enhances penetration of FD-4 kDa through zebrafish epidermis. As cCPE_{194–319}

could be a valuable Cldnb modulator for analysis of epithelial barrier functions, it is worthy to study whether cCPE_{194–319} might be used as a promising transepithelial enhancer during drug delivery.

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Author contributions

J.Z., C.N., Y.F. and J.P. designed the research; J.Z., C.N., Z.Y., A.P., H.C. and S.W. performed the research; A.P., Z.Q. and J.P. contributed new reagents/analytic tools; J.Z., C.N., Z.Y., Y.F. and J.P. analysed the data; and J.Z., C.N. and J.P. wrote the paper.

Conflict of interests

The authors have declared no conflicting interests.

Supporting Information

Additional supporting data may be found in the supplementary information of this article.

Figure S1. Inhibition of Cldnb causes developmental defects on zebrafish embryos at 48 hpf.

Figure S2. GST-cCPE_{194–319} protein binds to Cldnb in the epidermal cell membrane of zebrafish embryos.

Figure S3. The defective morphology caused by the absence of Cldnb is maintained at 72 hpf.

Figure S4. GST-cCPE_{194–319} protein treatment impairs epidermal barrier function in mice.

Table S1. Zebrafish heart beating rates after cCPE_{194–319} treatment with or without recovery.

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